P E SUBCLASS

NPR D TORRES

AIM-LOI

Serial No.: 09/763,957 Docket No.: 229752001300

Inventor: Jose Ramon Botella MESA et al.

Title: A PLANT PROMOTER AND USES THEREFOR

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AIM-I OLIGONUCLEOTIDES

-Oligonucleotide primers (SEQ ID NOS:4-9) used during Long Distance Inverse PCR

-Oligo's bind to regions of AIM-I (Mung bean ACC Synthase)

NSE-1 to NSE-6 correspond to SEQ ID NOS:4-9, respectively.

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NSE-1

5' -GCGGAT↓CCATCTTGGACAACAAGGGAGTT - 3'

29'omer

Tm = 68

NSE-2

5' -TAGGATC↓CAGAAAGACACTGAGAACCGTGG-_3'

30'omer

Tm = 70

NSE-3

5' -ACGGATCC↓GGTGTATGTGGTTAGAGTGTG- 3'

29'omer

Tm = 62

NSE-4

5' -CAGGATC↓CAGACATAGAGTGTGACCGCAA- 3'

29'omer

Tm = 66

NSE-5

5' -ATCGATCATATGAGCTCTAGACCCGGGCTGCAGGATCC ↓ GGTGTATGTGGTFAGAGTGTG- 3' 59'omer

Tm = 62

note: NSE-5 is identical to NSE-3 except different restriction enzyme sites have been incorporated (ie. 5'-Cla I, Nde I, Sac I, Xba I, Sma I, Pst I & Barn HI-3')

NSE-6

5' -CCGCGGAGATCTATCGATCTCGAGAATTC<u>AAGCTT</u> ↓ CAGACATAGAGTGTGACCGCAA-3' 57'omer

Tm = 66

note: NSE-6 is identical to NSE-4 except different restriction enzyme sites have been incorporated (ie. 5'-Sac II, Bgi II, CIa I, Xho I, Eco RI, & Hind III-3')

FIG. SUBCLASS APR 0 4 2003

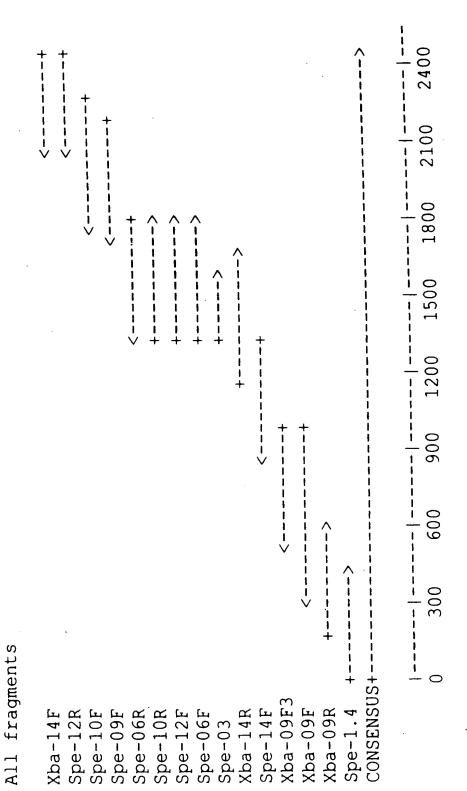
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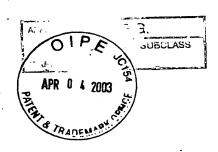
Inventor: Rose Ramon Botella MESA et al.
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FIGURE 4(i)

FIGURE 4(ii)

Figure 4 corresponds to SEQ ID NO:3

FIGURE 4

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GEL-1 2.5Kb Promoter Fragment

Len

Length: 2	2470	i uginione			
Length. 2	TTACAGATAC	ACAGAATCAG	ACGACACATC	TACTTTAATA	ACAGAAAAAT
51	AATAAGTGTC	GGAGATTATG	GTACGACAAG	ATGAAATGTT	TTTATATGGT
101	TGAGATTATT	TTGGTCTGTT	GTTGGAAGTT	TCACGAATCA	TGATTTTGAT
151	TTTACGTATT	AAAAAATGAA	AAGTTGAATC	ATGCATTTTA	TCTAGAAGCT
201	GGGAACTGAA	ССАААААААТ	AGCCAGTTGA	ACAACTGCAG	TATTTGTAGG
251	CGTATTCATT	TCTCCTTTCC	TACAATAATC	CTTGGTTGCT	CTTTATCGGA
301	ААААААССАА	AAGCAATAGC	TACTCTGTAA	GGTCCTCGAT	TGCCGACAAG
351	AACATCACAT	GCGTGCTGTC	GAAGAACACA	TAATTTTGAG	GTTGAAGCTC
401	ACGTGCGAGT	TTTGCATATT	TTTAGGTTAT	GTGTACACGT	ATGGAGTGAG
451	TTCCGCGTAT	ATAGTGTAGG	TAGTTGAGTG	GCTGAGTAGC	GAGTGAATCA
501	GGTAACACTA	TCTTTTCAAG	CCACCTAATT	AAGGGATTTA	ATGTTCATGC
551	AACTGTTCTT	CGCTAACTAA	GGCCCCACTT	ACCTTTATAA	TATTCTCTCT
601	AACTCCGGGC	TTTTGGTAAG	TACAACTTTT	CTACTCTTAT	TTAATGGAGG
651	GATTATTTTT	TCCATATACC	AATTAATTTA	TTTTTTAATT	TATGCATTTT
701	GATCTTATAT	TAAAACAATT	ATGGTATGGA	TTAAGTCGTA	TATCGGTGAC
751	AATTGAAGTT	TTCCTCAAGT	TTAGCCATTT	TTATGAAATT	AAACTTAATC
801	ACTACTATTA	GGTAAATTCA	TATGTATCAT	TAACAATTTC	AATGTGAGTT
851	CAATTTTACC	CAAGATTTGA	AAGTTGTTGT	CAACTTCTGT	TAACTAAAGT
901	TGTATTATAA	GGTTGACGAC	TTTAACCTAA	ATCTATTTTG	AATTGAAGGG
951	GTTGATGACT	TCAGCTTTAA	AATAATTCAA	CTAAAGTTCT	AGACTACATT
1001	GGAGATTTTA	GTGTTCATAA	AATTTTAGAA	AAAGGCTGAG	ТТАААСТТАТ
1051	GAAAAAGATT	GGTGACTATT	CAATTAATTA	GTTGTGAATT	GATGACAAAT
1101	ATTTCATGAG	CATAACCAAT	CAGAGAAATA	CCACCTCGAC	CGACTACAAC
1151	AATCTCAATG	TTAATTAATG	AAGCATTGTA	GTATAAGGAG	TCTAGAATAA
1201	ATTTCTTAAA	TATTAGAGGA	AAACTATTTT	ТАААААТТА	CAAGAAAAGT
1251	TTGATCTATA			ATCTAACAAT	TTTCTTATGA
FIGURE 4(i)					





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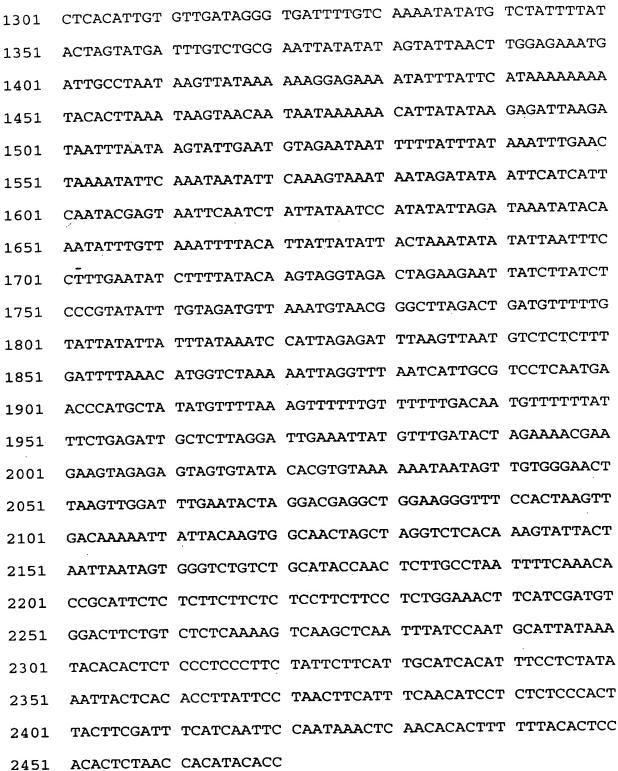
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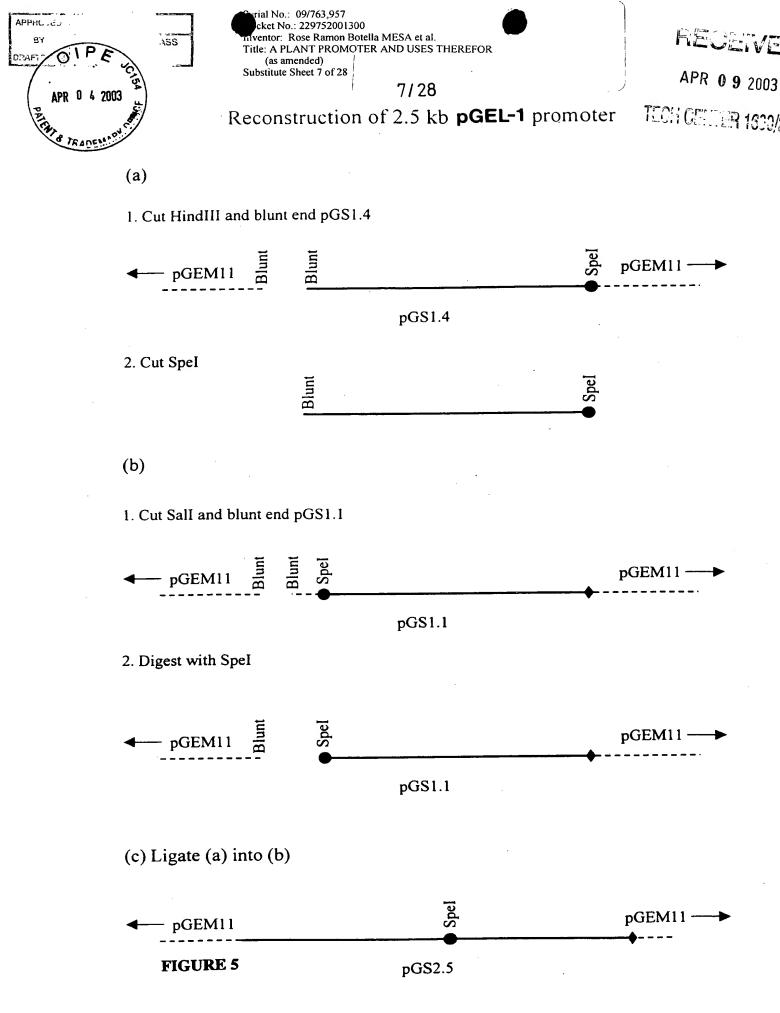


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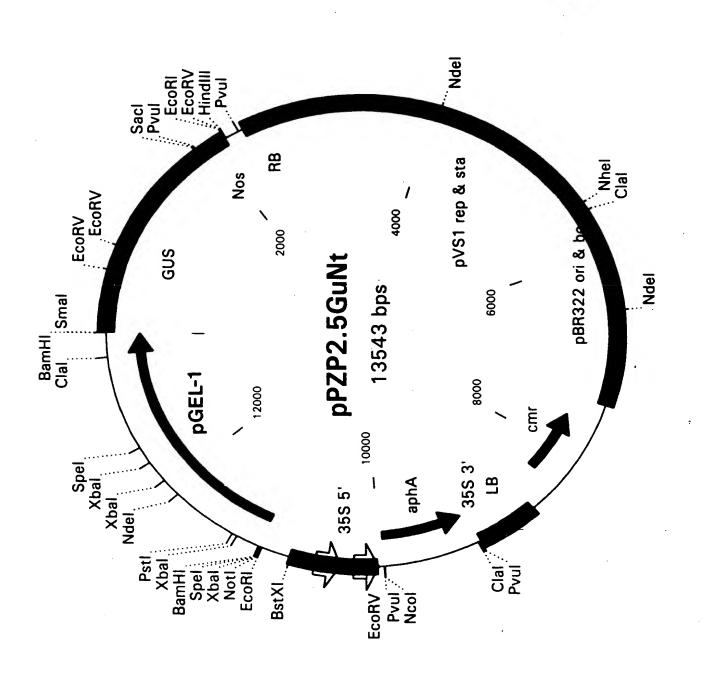
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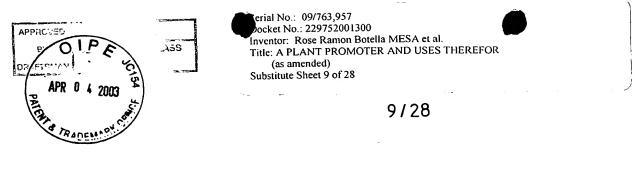
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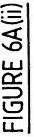
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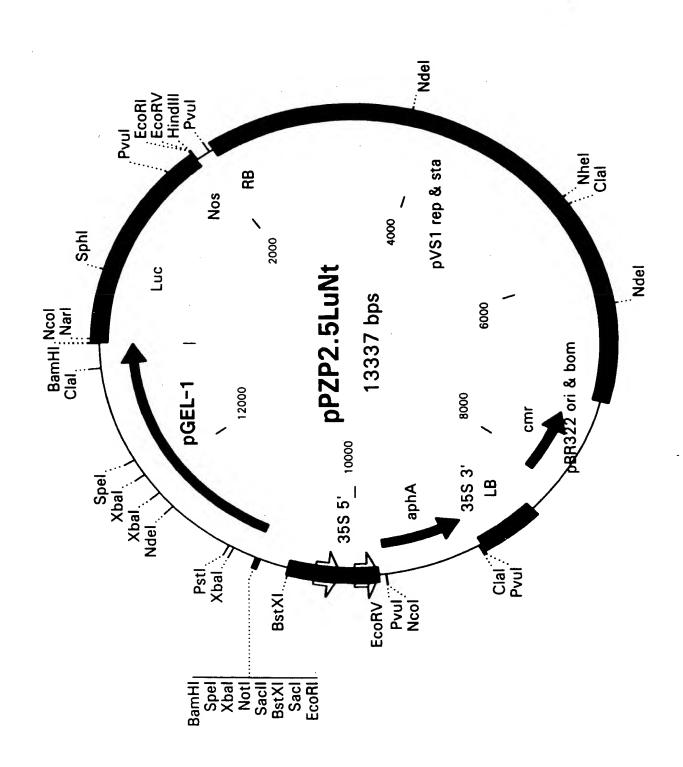
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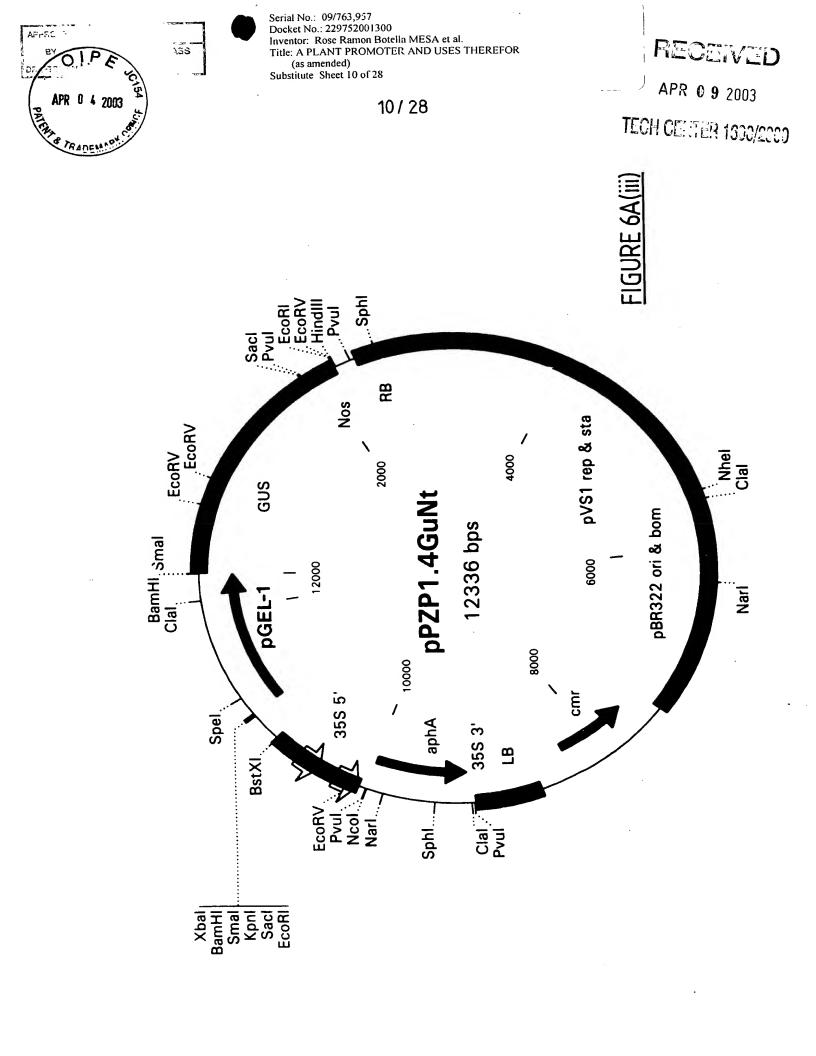
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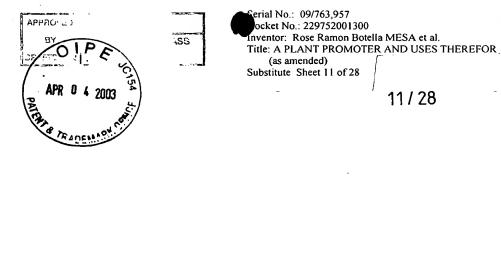










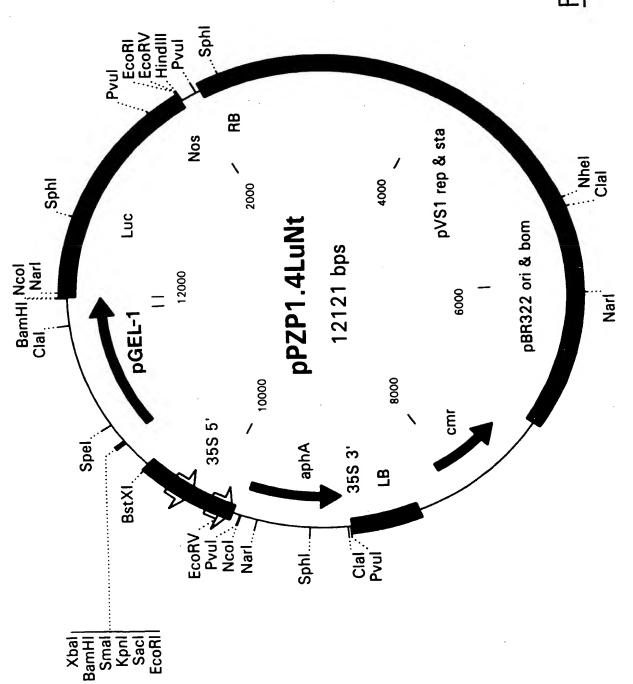


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FIGURE 6A(iv)





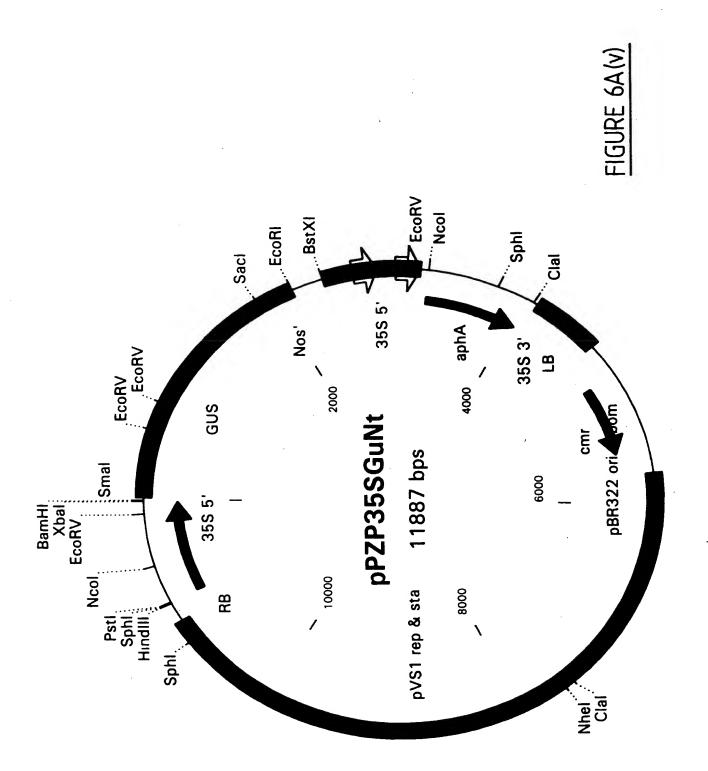
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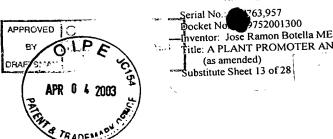
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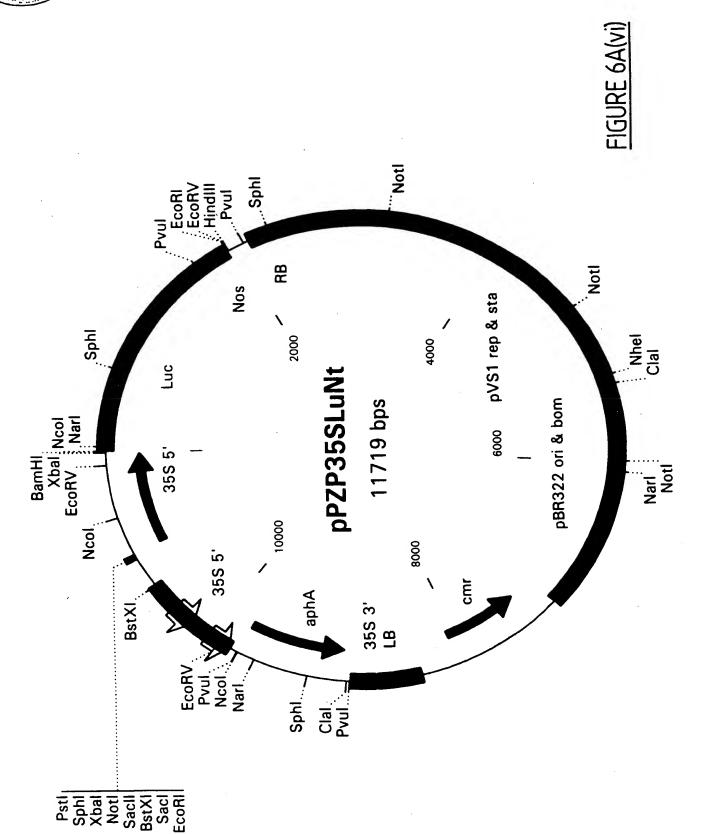
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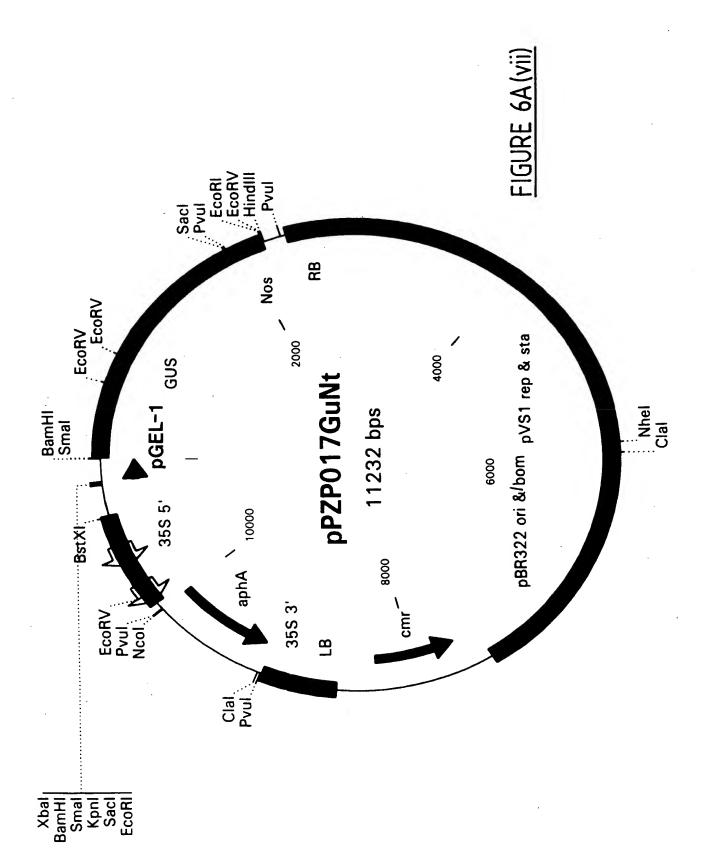
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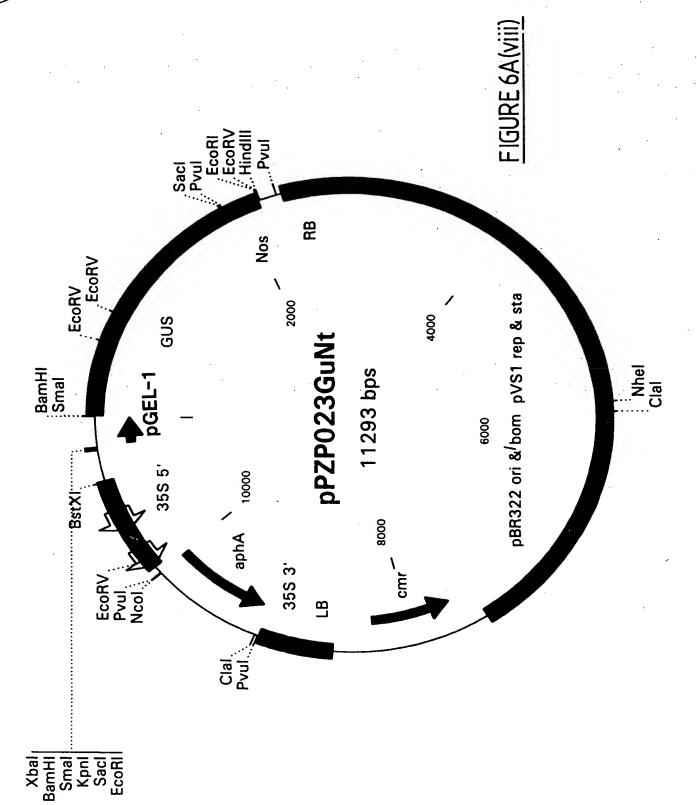
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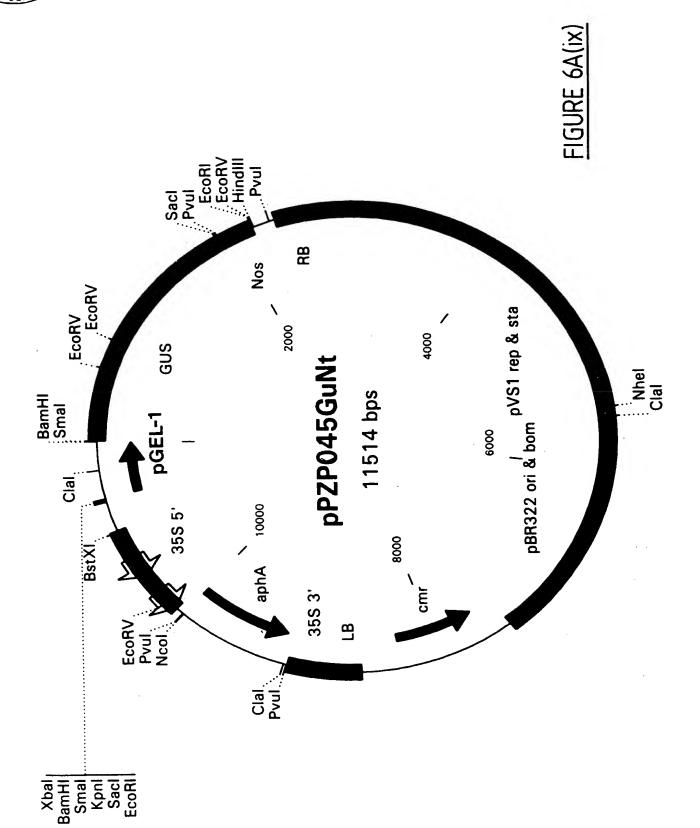
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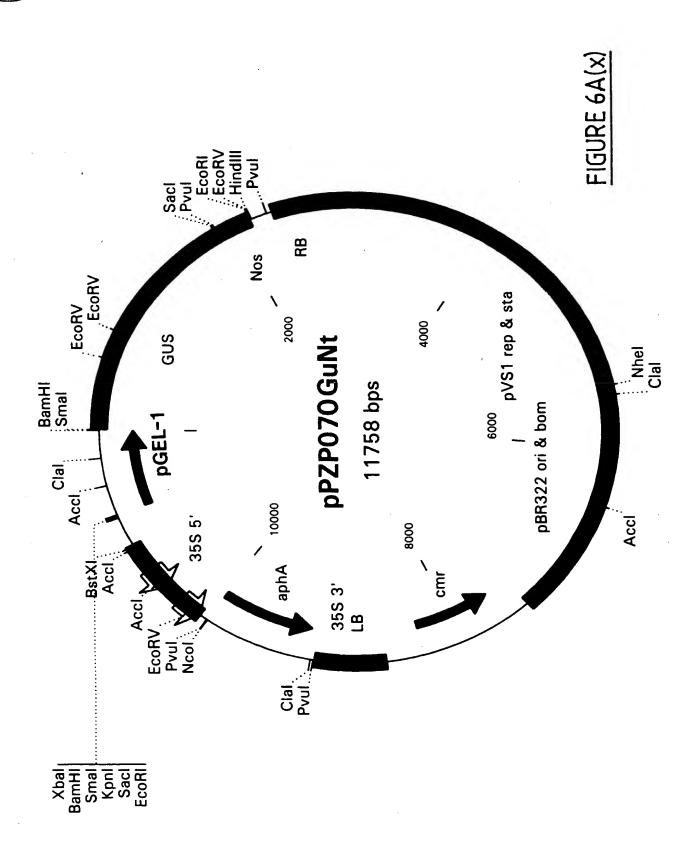
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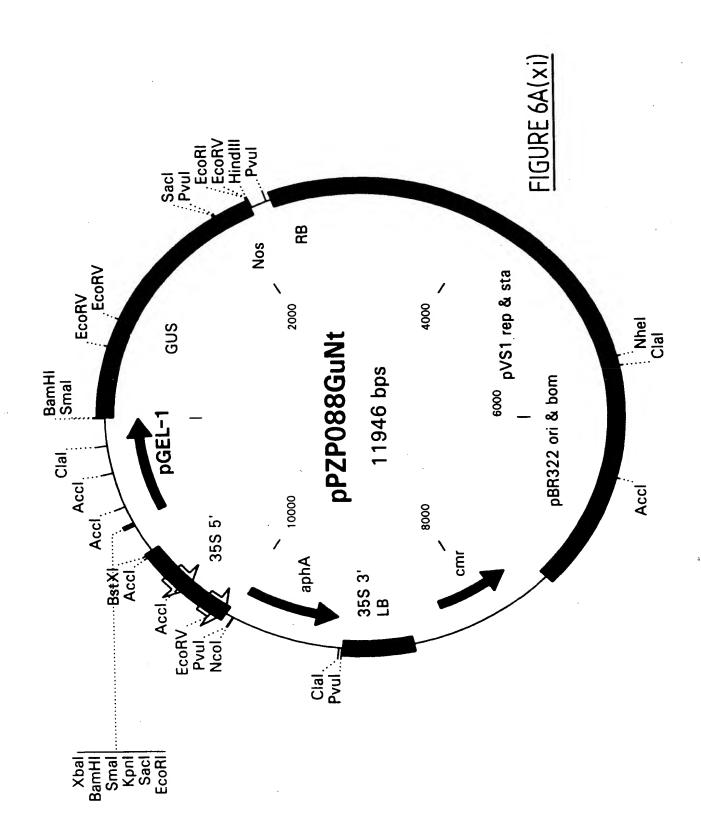
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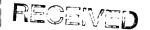
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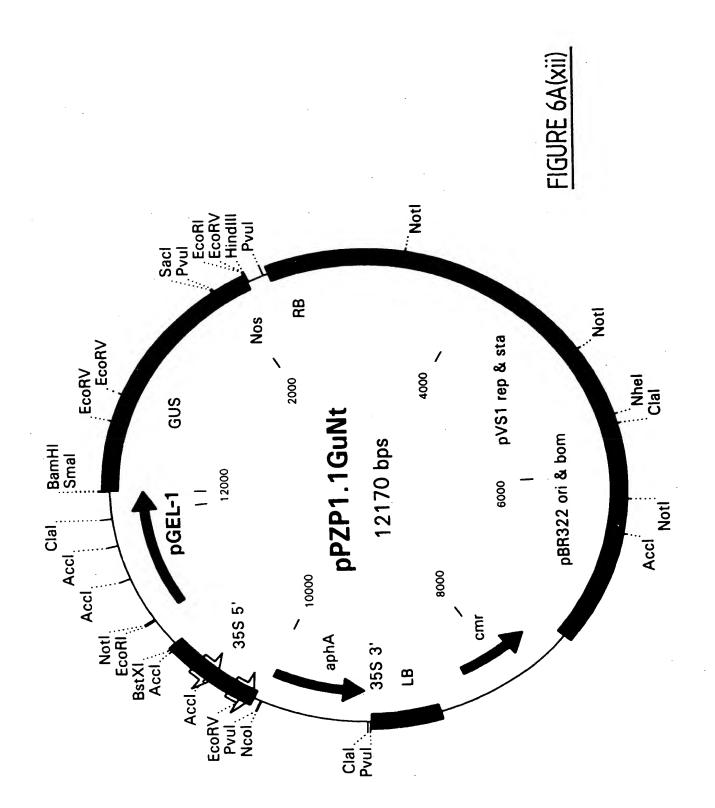
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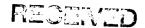




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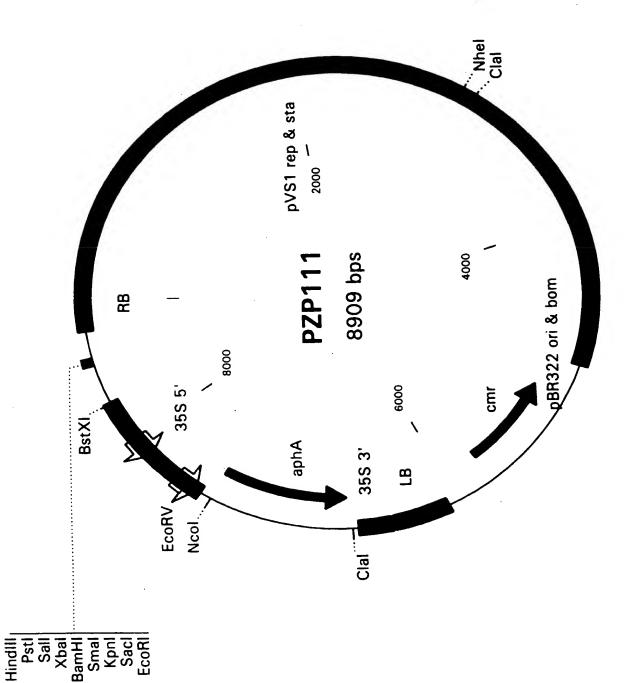
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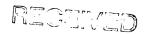
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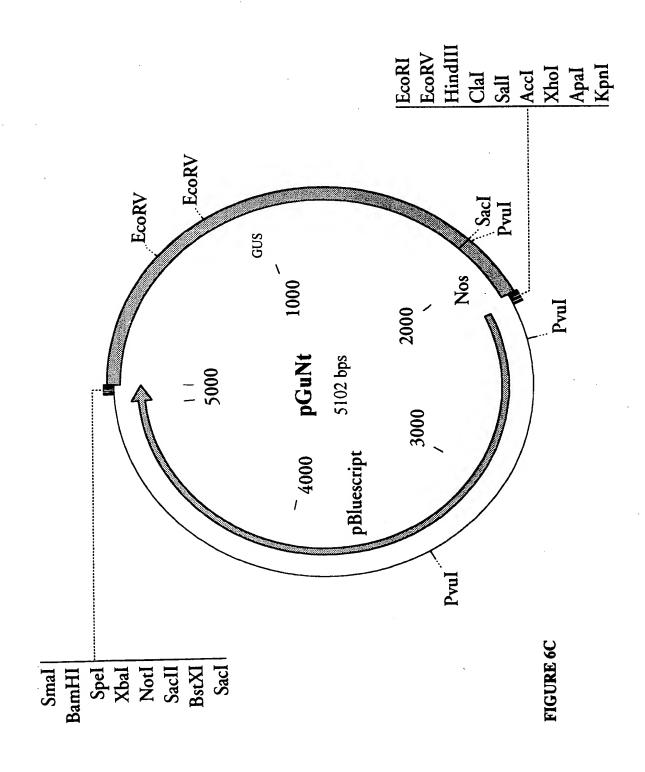
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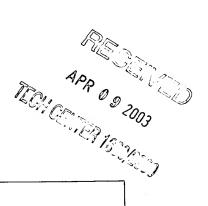


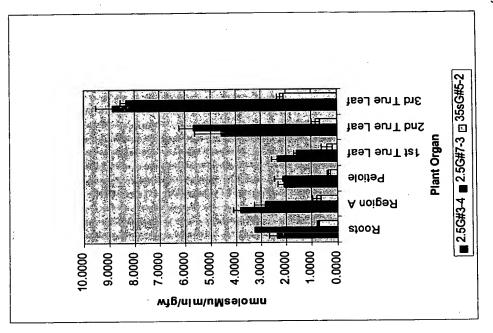
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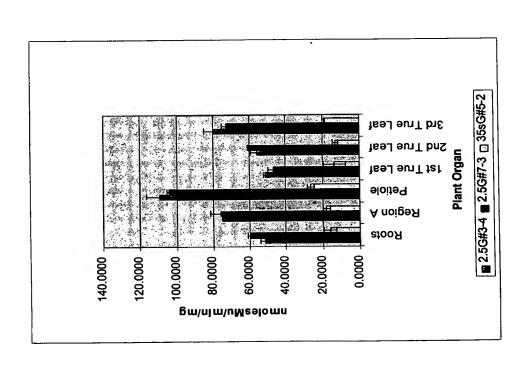
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IGURE 8B



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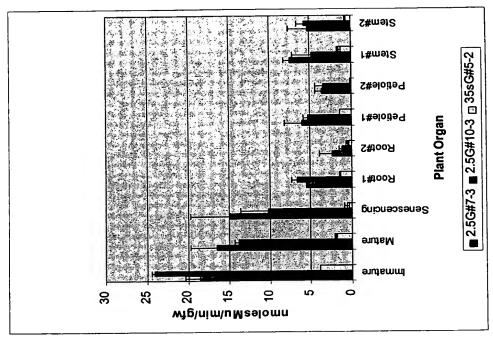
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IGURE 9B

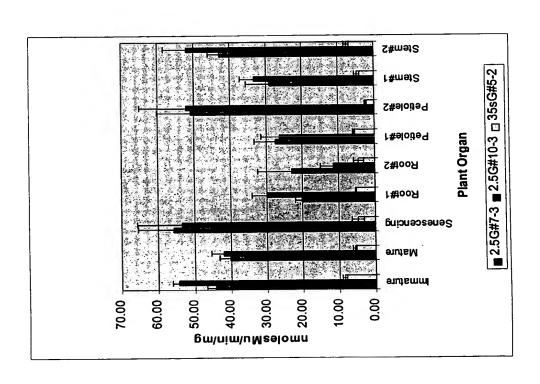
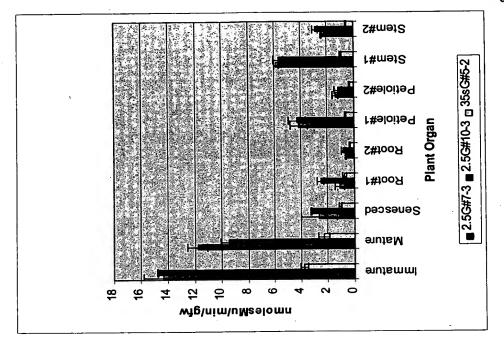


FIGURE 94

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IGURE 10B

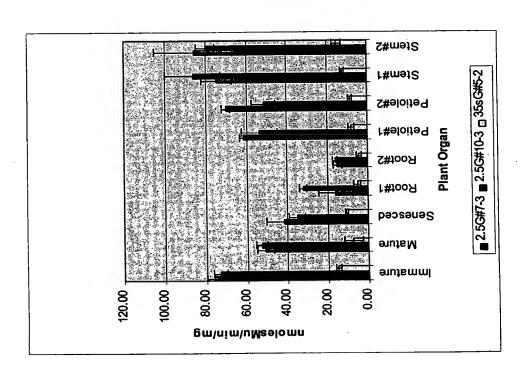


FIGURE 10A



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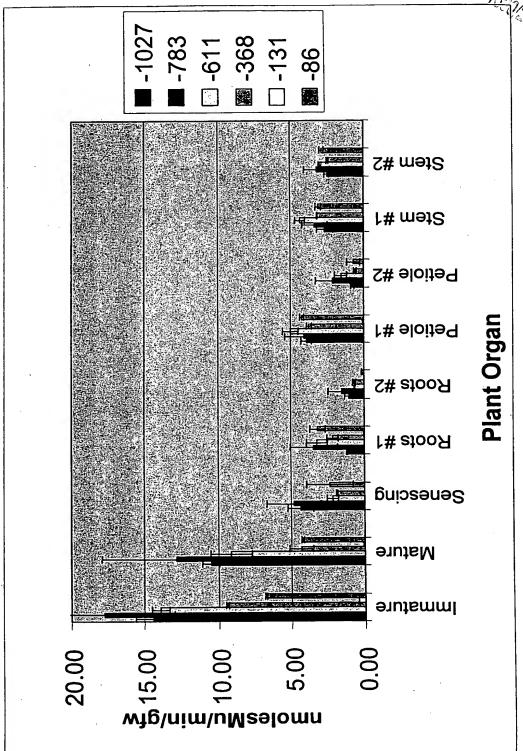
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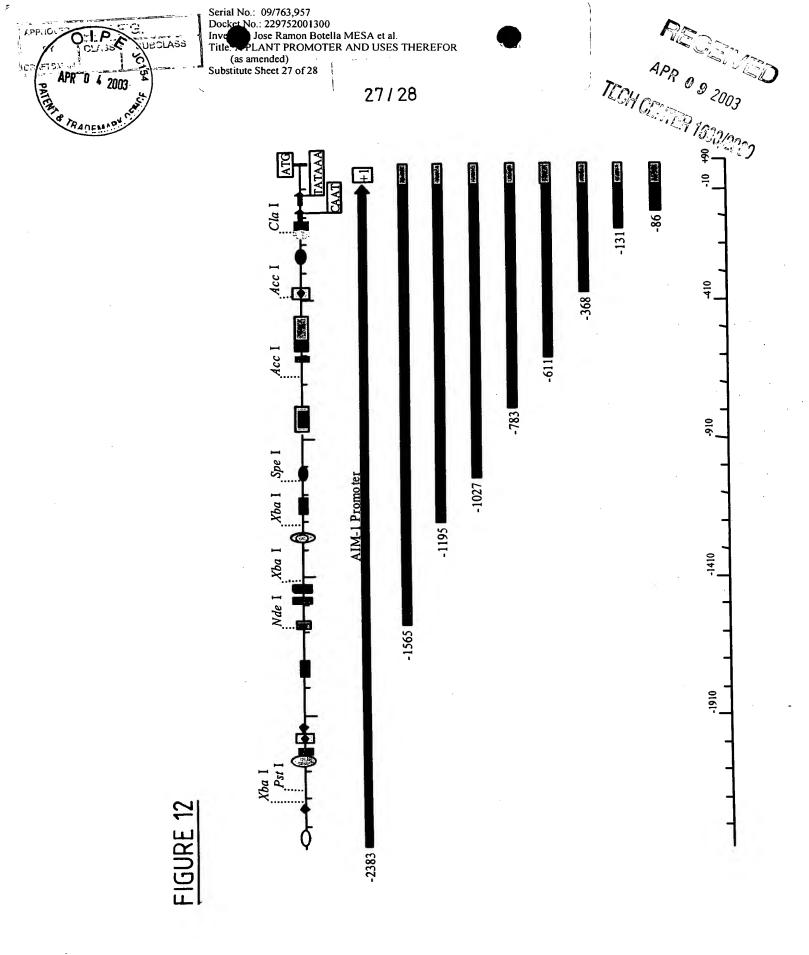
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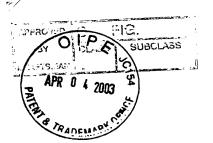
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IGURE 11





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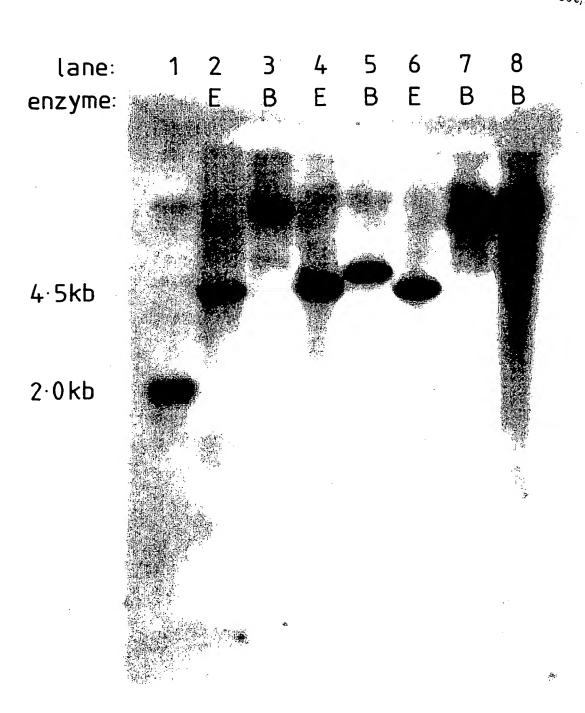


FIGURE 13